**Details**

1. **Chlamy\_TF\_Family:** First TF-gene name list of around 200 gens
2. **ExpCCM2019csv-Copy:** CCM-gene name list of around 100 genes, from 2019 paper
3. **Normalized\_counts\_chlamy\_EdgeR\_2019\_data:** Expression data of all genes from three time-series experiments, from 2019 paper
4. **expall48\_ccm\_tf\_fromR:** Normalized\_counts\_chlamy\_EdgeR\_2019\_data, filtered by the Cre ID of the genes that are either CCM genes or TF genes (processed data using 1-3 by R (no code attached))
5. **updated\_TF\_list\_Citu\_June2k20**: Second TF-gene name list of around 700 gens
6. (**ErrorListGene:** Genes that throw out errors in dynGENIE3)
7. (**Gene\_names:** the pairs of gene number and Cre ID)